Fast Sequence Lookup


```
LOCUS       AY008270                562 bp    DNA     linear   PRI 07-OCT-2000
DEFINITION  Homo sapiens cholesteryl ester transfer protein (CETP) gene, intron
            12 and partial cds.
ACCESSION   AY008270
VERSION     AY008270.1  GI:10716515
KEYWORDS    .
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 562)
  AUTHORS   Travali,S., Emmanuele,G., Cefalu,A.B., Noto,D., Marino,G.,
            Avondo,S., Litrico,C. and Averna,M.R.
  TITLE     Characterization of CETP intron 12 sequence disparity with
            published sequences
  JOURNAL   Unpublished
REFERENCE   2  (bases 1 to 562)
  AUTHORS   Travali,S., Emmanuele,G., Avondo,S. and Litrico,C.
  TITLE     Direct Submission
  JOURNAL   Submitted (20-SEP-2000) Dept. Biomedical Sciences, University of
            Catania, Via Androne 87, Catania, CT 95124, Italy
REFERENCE   3  (bases 1 to 562)
  AUTHORS   Cefalu,A.B., Noto,D., Marino,G. and Averna,M.R.
  TITLE     Direct Submission
  JOURNAL   Submitted (03-OCT-2000) Dept. Internal Medicine and Geriatrics,
            University of Palermo, Via de Vespro 141, Palermo, PA 90127, Italy
FEATURES             Location/Qualifiers
     source          1..562
                     /organism="Homo sapiens"
                     /mol_type="genomic DNA"
                     /db_xref="taxon:9606"
     gene            <1..>562
                     /gene="CETP"
     intron          <1..14
                     /gene="CETP"
                     /number=11
     mRNA            join(<15..82,492..>525)
                     /gene="CETP"
                     /product="cholesteryl ester transfer protein"
     CDS             join(<15..82,492..>525)
                     /gene="CETP"
                     /codon_start=1
                     /product="cholesteryl ester transfer protein"
                     /db_xref="/protein_id="AAG21822.1"
                     /db_xref="GI:10716516"
                     /translation="DIVTTVQASYSKKKLFLSLLDFQITPKTVSNLTE"
     exon            15..82
                     /gene="CETP"
                     /number=12
     intron          83..491
                     /gene="CETP"
                     /number=12
```

```
     exon            492..525
                     /gene="CETP"
                     /number=13
     intron          526..>562
                     /gene="CETP"
                     /number=13
ORIGIN
        1 gtttctctcc ccaggatatc gtgactaccg tccaggcctc ctattctaag aaaaagctct
       61 tcttaagcct cttggatttc cagtatgtgc tgcagagaag agaggggcg gtcaactccg
      121 caaacctctc cctggcccct tggagtcagg cacagggcgg ggtgttggtg gggaaatgtg
      181 gccccttttct tctggggcat atgggctgac tgcagggaga taagaccctg cctagataga
      241 atcttcgtgg ggaagaaggg gctccaggaa gaatggaggg agggcctggc aggaggagag
      301 cgctgcccga gcaaaggcct ggccgccaga atagcaaatc tcaagggaat agcaaatctc
      361 aagagagtgc cccaaagggc ctgagctatg agacaaaagc actggctgct attcttagag
      421 tttctttccc aggggatgtt acaggagggg gcccaatgga gggtcaaatt atcatcgctt
      481 ttttatttca ggattacacc aaagactgtt tccaacttga ctgaggtagg tagtcttgga
      541 tagactgggg gaaataagtc ct
//
```
